# Mutants-P318A.ST25 SEQUENCE LISTING

	SEQUENCE LISTING	
<110>	Covalys Biosciences AG	
<120>	Mutants of O6-Alkylguanine-DNA Alkyltransferase	
<130>	P318A	
<150> <151>	EP04405123.3 2004-03-02	
<150> <151>	EP04405465.8 2004-07-22	
<160>	48	
<170>	PatentIn version 3.3	
<210> <211> <212> <213>	1 624 DNA Homo sapiens	
<400>	1 aagg attgtgaaat gaaacgcacc acactggaca gccctttggg gaagctggag	60
		120
	99 9-9-9	180
	ggggggggggggg	240
		300
	guar cource cg	360
		420
		480
		540
		600
		624
J		
<210> <211> <212> <213>	2 22 DNA Artificial Sequence	
<220>		
<223>	Substrate oligonucleotide containing O6-Benzylguanine at posit 14	ion
<220> <221> <222> <223>	misc_feature (14)(14) n is O6-benzylguanine	
<400> gtggtg	2 ggca gctnaggcgt gg	22
<210> <211> <212> <213>	3 33 DNA Artificial Sequence	

	Mutants-P318A.ST25	
<220> <223>	Sense primer for cloning AGT into pGEX	
<400> cgaaat	3 ggat ccatggacaa ggattgtgaa atg	33
<210> <211> <212> <213>	4 43 DNA Artificial Sequence	
<220> <223>	Antisense primer for cloning AGT into pGEX	
<400> gccttt	4 gaat tccgtctttg tagtcgtttc ggccagcagg cgg	43
<210> <211> <212> <213>	5 31 DNA Artificial Sequence	
<220> <223>	Sense primer for mutating K125A, T127A, R128A	
<400> gcaacc	5 ccgc agccacggca gcagtgggag g	31
<210> <211> <212> <213>	6 31 DNA Artificial Sequence	
<220> <223>	Antisense primer for mutating K125A, T127A, R128A	- 47-4
<400> cctccc	6 actg ctgccgtggc tgcggggttg c	31
<210> <211> <212> <213>	7 49 DNA Artificial Sequence	
<220> <223>	Sense primer for cloning into eukaryotic pNUC vector	
<400> gatcga	7 gcta gcgctaccgg tcgccaccat ggacaaggat tgtgaaatg	49
<210> <211> <212> <213>	8 31 DNA Artificial Sequence	
<220> <223>	Antisense primer for cloning into eukaryotic pNUC vector	
<400> gctagg	8 gatc ctacgtttcg gccagcaggc g	31
<210> <211> <212> <213>	9 35 DNA Artificial Sequence	

### Mutants-P318A.ST25

```
<220>
<223>
       Sense primer for mutating Cys 62 to Ala
<400>
                                                                        35
gagcccctga tgcaggctac agcctggctg aatgc
<210>
       10
<211> 35
<212> DNA
<213> Artificial Sequence
<220>
<223> Antisense primer for mutating Cys 62 to Ala
<400> 10
gcattcagcc aggctgtagc ctgcatcagg ggctc
                                                                        35
<210>
       11
<211> 60
<212> DNA
<213> Artificial Sequence
<220>
       Sense primer for cloning of AGT mutants into phage-display vector
<223>
<400> 11
ctactcgcgg cccagccggc catggcggac tacaaagaca tggacaagga ttgtgaaatg
                                                                        60
<210>
       12
<211>
       40
<212>
       DNA
      Artificial Sequence
<213>
<220>
<223>
       Antisense primer for cloning of AGT mutants into phage-display
<400> 12
                                                                        40
ggaattcggc ccccgaggcc gcgtttcggc cagcaggcgg
<210>
<211>
       42
<212> DNA
<213> Artificial Sequence
<220>
<223>
      Antisense primer for cloning AGT truncated after 182 into pGEX
<400> 13
gcctttgaat tccgtctttg tagtctccca agcctggctt cc
                                                                        42
<210>
       14
<211>
       50
<212>
      DNA
<213> Artificial Sequence
<220>
       Sense primer for randomisation of codons 131-135
<220>
<221> misc_feature
<222> (22)..(23)
<223> n is a, c, g, or t
```

### Mutants-P318A.ST25 <220> misc\_feature <221> (25)..(26)<222> n is a, c, g, or t <223> <220> <221> misc\_feature <222> (31)..(32)n is a, c, g, or t <223> <220> misc\_feature <221> <222> (34)..(35)<223> n is a, c, g, or t 50 cccaaagccg cgcgagcagt gnnknnkgca nnknnkggca atcctgtccc <210> 15 <211> 26 <212> DNA <213> Artificial Sequence <212> <220> Antisense primer for randomisation of codons 131-135 <223> <400> 15 26 tgctcgcgcg gctttggggt tgcctg <210> 16 <211> 41 <212> DNA <213> Artificial Sequence <220> Sense primer for randomisation of codons 115-116 <223> 213 <220> misc\_feature (19)..(20) <221> <222> <223> n is a, c, g, or t <220> <221> <222> misc\_feature (22)..(23) n is a, c, g, or t <223> 41 ggagaagtga tttcttacnn bnnbttagca gccctggcag g <210> <211> 27 <212> DNA <213> Artificial Sequence <223> Antisense primer for randomisation of codons 115-116 <400> 17 27 gtaagaaatc acttctccga atttcac <210> 18 <211> 41 <212> DNA <213> Artificial Sequence

## Mutants-P318A.ST25 <220> <223> Sense primer for randomisation of codons 150-152 <220> <221> misc\_featu <222> (19)..(20) misc\_feature <223> n is a, c, g, or t <220> <221> misc\_feature <222> (22)..(23) <223> n is a, c, g, or t <220> <221> misc\_feature <222> (25)..(26) <223> n is a, c, g, or t ccgtgccaca gagtggtcnn bnnbnnbgga gccgtgggcg g 41 <210> 19 <211> <212> 18 DNA <213> Artificial Sequence <223> Antisense primer for randomisation of codons 150-152 <400> 19 18 gaccactctg tggcacgg <210> 20 <211> 35 <212> DNA <213> Artificial Sequence <220> <223> Sense primer for mutating G131K, G132T, M134L, R135S <400> 20 35 gcagccacgg cagcagtgaa gacggcactg agtgg <210> 21 <211> 40 <212> DNA <213> Artificial Sequence <220> <223> Antisense primer for mutating G131K, G132T, M134L, R135S <400> 21 40 ggatagggac aggattgcca ctcagtgccg tcttcactgc <210> 41 <211> <212> DNA <213> Artificial Sequence <220> <223> Sense primer for mutating Q115S, Q116H <400> 22 41 gtgaaattcg gagaagtgat ttcttactct cacttagcag c

	Mutants-P318A.ST25	
<210> <211> <212> <213>	23 35 DNA Artificial Sequence	
<220> <223>	Antisense primer for mutating Q115S, Q116H	
<400> cctgcc	23 aggg ctgctaagtg agagtaagaa atcac	35
<210> <211> <212> <213>	24 33 DNA Artificial Sequence	
<220> <223>	Sense primer for mutating C150N, S151I, S152N	
<400> cgtgcc	24 acag agtggtcaat atcaatggag ccg	33
<210> <211> <212> <213>	25 32 DNA Artificial Sequence	
<220> <223>	Antisense primer for mutating C150N, S151I, S152N	
<400> cgtaac	25 cgcc cacggctcca ttgatattga cc	32
<210> <211> <212> <213>	26 32 DNA A Artificial sequence	
<220> <223>	Primer for cloning mutant AGT in pET15b	
<400> gtcgca	26 tatg gacaaggatt gtgaaatgaa ac	32
<210> <211> <212> <213>	27 34 DNA Artificial sequence	
<220> <223>	Primer for cloning mutant AGT in pET15b	
<400> gattac	27 ggga tccttatccc aagcctggct tccc	34
<210> <211> <212> <213>	28 45 DNA Artificial Sequence	
<220> <223>	Antisense primer for cloning truncated AGT in pAK 100	
<400> gcaatg	28 gaat tcggcccccg aggccgctcc caagcctggc ttccc	45

## Mutants-P318A.ST25

<210> <211> <212> <213>	29 81 DNA Artificial Sequence	
<220> <223>	Sense primer for introducing 12xHis, SbfI and AscI sites	
<400> ttatcc	29 atgg cacatcatca tcatcatcat catcatcatc atcatcatcc tgcaggtata	60
ggcgcg	ccta aaagcttctt a	81
<210> <211> <212> <213>	30 81 DNA Artificial Sequence	
<220> <223>	Antisense primer for introducing 12xHis, SbfI and AscI sites	
<400> taagaa	30 gctt ttaggcgcgc ctatacctgc aggatgatga tgatgatgat gatgatgatg	60
atgatg	atgt gccatggata a	81
<210> <211> <212> <213>	31 59 DNA Artificial Sequence	
<220> <223>	Sense primer for cloning mutant AGT in pBAD-HisA	
<400> ggcctg	31 cagg tgaaaacctg tacttccagg gtatggacaa ggattgtgaa atgaaacgc	59
<210> <211> <212> <213>	32 45. DNA Artificial Sequence	
<220> <223>	Antisense primer for cloning mutant AGT in pBAD-HisA	
<400> aaaagg	32 cgcg ccggatcctt atcccaagcc tggcttcccc aaccg	45
<210> <211> <212> <213>	33 47 DNA Artificial Sequence	
<220> <223>	Antisense primer for cloning wt AGT in pBAD-HisA	
<400> aacggc	33 gcgc cggatcctta gtttcggcca gcaggcgggg agcccga	47
<210> <211> <212> <213>	34 49 DNA Artificial Sequence	
<220>		

<223>	Mutants-P318A.ST25 Sense primer for cloning AGTM in pEGFP-Nuc		
<400> gatcga	34 gcta gcgctaccgg tcgccaccat ggacaaggat tgtgaaatg		49
<210> <211> <212> <213>	35 32 DNA Artificial Sequence		
<220> <223>	Antisense primer for cloning AGT G160W in pEGFP-Nuc		
<400> ccaggc	35 agat ctgtttcggc cagcaggcgg gg		32
<210> <211> <212> <213>	36 48 DNA Artificial Sequence		
<220> <223>	Antisense primer for cloning AGTM in pEGFP-Nuc		
<400> ccaggc	36 agat cttcccaagc ctggcttccc caaccggtgg ccttcatg		48
<210> <211> <212> <213>	37 33 DNA Artificial Sequence		
<220> <223>	Sense primer for cloning beta-Gal in pEGFP-Nuc		
<400> catcgt	370 ctag attattttg acaccagacc aac	.a	33
<210> <211> <212> <213>	38 42 DNA Artificial Sequence		
<220> <223>	Antisense primer for cloning beta-Gal in pEGPF-Nuc		
<400> gatcga	38 gatc tgggtccgga atgactaaat ctcattcaga ag		42
<210> <211> <212> <213>	39 25 DNA Artificial Sequence		
<220> <223>	Sense primer for mutation G160W		
<400> caacta	39 ctcc tggggactgg ccgtg		25
<210><211><211><212><213>	40 25 DNA Artificial Seguence		

220	Mutants-P318A.ST25	
<220> <223>	Antisense primer for mutation G160W	
<400> ccagto	40 ccca ggagtagttg cccac	25
<210> <211> <212> <213>	41 39 DNA Artificial Sequence	
<220> <223>	Primer for error prone PCR of pAK $100$ insert, c at position 1 modified with biotin	
<220> <221> <222> <223>	misc_feature (1)(1) n is c modified with biotin	
<400> ngatco	41 ttag acctgaacgc aggtttcccg actggaaag	39
<210> <211> <212> <213>	42 39 DNA Artificial Sequence	
<220> <223>	Antisense primer for error prone PCR of pAK100 insert	
<400> gcgtca	42 gggt tacaagttca tggtttacca gcgccaaag	39
<210> <211> <212> <213>		
<220> <223>	Sense primer for amplification of errorprone-PCR product	
<400> cgatcc	43 ttag acctgaacg	19
<210> <211> <212> <213>	44 19 DNA Artificial Sequence	
<220> <223>	Antisense primer for amplification of errorprone-PCR product	
<400> gcgtca	44 gggt tacaagttc	19
<210> <211> <212> <213>	45 48 DNA Artificial Sequence	
<220> <223>	Sense primer for saturation mutagenesis AGTM 150-154	

```
Mutants-P318A.ST25
<220>
<221> misc_feature
<222> (18)..(19)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (21)..(22)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (24)..(25)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222>
        (27)..(28)
<223> n is a, c, g, or t
<220>
<221> misc_feature <222> (30)..(31)
<223> n is a, c, g, or t
<400> 45
                                                                                       48
cgtgccacag agtggtcnnk nnknnknnkn nkgtgggcgg ttacgagg
<210> 46
<211> 48
<212> DNA
<213> Artificial Sequence
<223> Antisense primer for saturation mutagenesis AGTM 150-154
<220>
                                                                             144
<221> misc_feature
<222> (18)..(19)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (21)..(22)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222>
        (24)..(25)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (27)..(28)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (30)..(31)
<223> n is a, c, g, or t
cctcgtaacc gcccacmnnm nnmnnmnnmn ngaccactct gtggcacg
                                                                                       48
<210> 47
<211> 48
<212> DNA
```

<213> Artificial Sequence

## Mutants-P318A.ST25

```
<220>
<223> Sense primer for saturation mutagenesis AGTM 31-35
<220>
<221>
       misc_feature
<222>
        (17) (18)
<223> n is a, c, g, or t
<220>
<221>
<222>
       misc_feature
(20)..(21)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (23)..(24)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (26)..(27)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (29)..(30)
<223> n is a, c, g, or t
<400>
gcagggtctg cacgaannkn nknnknnknn kaaggggacg tctgcagc
                                                                                     48
<210> 48
<211> 48
<212> DNA
<213> Artificial Sequence
<220>
<223> Antisense primer for saturation mutagenesis AGTM 31-35
<220>
<221>
<222>
       misc_feature
       (19)..(20)
<223> n is a, c, g, or t
<220>
<221>
<222>
        misc_feature
<222> (22)..(23)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222>
       (25)..(26)
<223> n is a, c, g, or t
<220>
<221> misc_feature
<222> (28)..(29)
<223> n is a, c, g, or t
<220>
<221>
<222>
       misc_feature
(31)..(32)
<223>
       n is a, c, g, or t
gctgcagacg tccccttmnn mnnmnnmnnm nnttcgtgca gaccctgc
                                                                                     48
```